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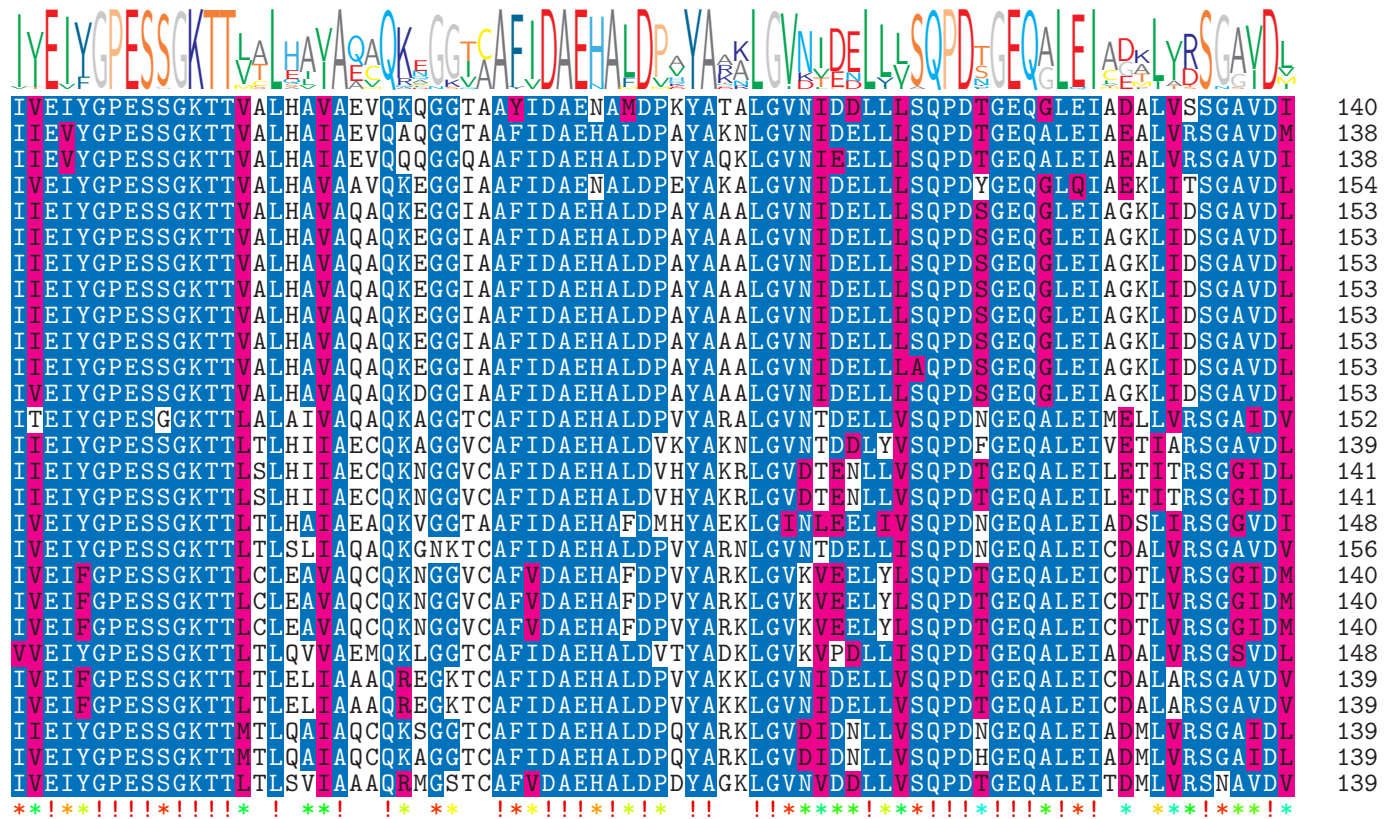
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NTDB id 269 KZH43 RS08665 WP 001085462.1
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NTDB id 1313 DR RS12030 WP 010888966.1
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NTDB id 1074 ABD1 RS10195 WP 000344167.1
NTDB id 1022 ACIAD RS06390 WP 004925639.1
NTDB id 1205 PSJM300 05545 AFN77183.1
consensus



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NTDB id 527 SMSK321 RS11065 WP 001085454.1	SVRLDVRGSTQIKGTG.DQKDTNVGKETKIKVVKNKVAPPFKAEAFVEIMYGGGISKTGELLKIASDLDIKKKAGAWYSYK	311
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consensus	*!!**!* * ** *****!!!!!!*! * * *****!*!*** ***** ** *** ! !*****!	

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NTDB id 1032760 AB8V66 RS04635 WP 003719674.1	EERIGGGRENAKQYLKEHTDIRDEISKRVREEYEIDGSNKEPL..AETEETLSLLDDE.....	348
NTDB id 115 BSU 16940 NP 389576.2	EERLGGGRENAKQFLKENKDIMLMIQEQUIREHYGLDNNGVVQQAEETQEELEFEE.....	348
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NTDB id 269 KZH43 RS08665 WP 001085462.1	DEKIGQGSSENAKKYLAHPPEIFDEIDKQVRSKFGGLIDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
NTDB id 228 SPD RS09265 WP 001085462.1	DEKIGQGSSENAKKYLAHPPEIFDEIDKQVRSKFGGLIDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
NTDB id 194 SPR RS08825 WP 001085462.1	DEKIGQGSSENAKKYLAHPPEIFDEIDKQVRSKFGGLIDGEEVSEQDTENKKDE....PKKEEA..VNEEVPLDLGDELEIE	385
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NTDB id 527 SMSK321 RS11065 WP 001085454.1	DEKIGQGSSENAKKYLADNPEVFAEIDHQVVRVQFGLIDGEEAAES....KKDE....VAQVDS..VNEEVTLDLGDELEIE	381
NTDB id 453 Spy49 1753c ACI62003.1	GEKIGQGSSENAKRYLADHPQLFDEIDRKVRVKFGLLEESEESAMAVASE.....E.....TDDLALDLNNGIEIE	377
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NTDB id 1142 GTF74 RS11550 WP 000344154.1	GDKIGQGGKANACKYLLKENPEIAKTLDKKLREMLLNPNM....QLIAETSSAADDVEFGA..VPEEF.....	354
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consensus	**!!*!* * * * * * * * * * *	

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NTDB id 625 LCA RS02525 WP 011374196.1	...	355
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NTDB id 1022 ACIAD RS06390 WP 004925639.1	...	349
NTDB id 1205 PSJM300 05545 AFN77183.1	...	347
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved